SEQUENCE LISTING														
<110> Dale, James Langham Echeverria, Santy Peraza														
<120> BANANA RESISTANCE GENES AND USES THEREOF														
<130> DAVI172.006APC														
<140> 10573372 <141> 2006-10-31														
<150> PCT/AU2004/001300 <151> 2004-09-23 <150> AU 2003905222 <151> 2003-09-25														
<151> 2004-09-23 <150> AU 2003905222 <151> 2003-09-25 <160> 7														
<150> AU 2003905222 <151> 2003-09-25 <160> 7														
<170> FastSEQ for Windows Version 4.0														
<170> FastSEQ for Windows Version 4.0 <210> 1 <211> 4380 <212> DNA														
<210- 1 <211- 4380 <212- DNA <213- Musa acuminata <220- <221- CDS <222- (1)(4323)														
<pre><400> 1 atg tcg acg gcg cta gta atc gga gga tgg ttc gcg caa agc ttc atc Met Ser Thr Ala Leu Val Ile Gly Gly Trp Phe Ala Gln Ser Phe Ile 1</pre>														
cag acg ttg ctc gac aag gcc agc aac tgc gcg atc caa caa ctc gcg Gln Thr Leu Leu Asp Lys Ala Ser Asn cys Ala Ile Gln Gln Leu Ala 20 25 30														
cgg cgc cgc cgc ctt cac gat gac ctg agg cgg ctg cgg acg tct ctg Arg Arg Arg Arg Gly Leu H1s Asp Leu Arg Arg Leu Arg Thr Ser Leu $^{\rm 448}$ $^{\rm 35}$														
ctc cgg atc cat gcc atc ctc gac aag gca gag acg agg tgg aac cat $$192$$ Leu Arg ITe His Ala ITe Leu Asp Lys Ala Glu Thr Arg Trp Asn His 50														
aaa aac acg agc ttg gtg gag ctg gtg agg cag ctc aag gat gct gcc Lys Asn Thr Ser Leu Val Glu Leu Val Arg Gln Leu Lys Asp Ala Ala 65 70														

336

caa aag gtc gag cac cgg gga gac cag ata agc gac ctc ttt tct ttt Gln Lys Val Glu His Arg Gly Asp Gln Tle Ser Asp Leu Phe Ser Phe 100 100 110

tcc Ser	ctt Leu	agt Ser 115	act Thr	gcg Ala	agc Ser	gag Glu	tgg Trp 120	ttg Leu	ggt	gcc Ala	gat	ggt	gat Asp	gat Asp	gct Ala	384
ggg Gly	act Thr 130	cga Arg	ttg Leu	agg Arg	gag Glu	atc Ile 135	cag Gln	ggg G1y	aag Lys	ctg Leu	tgc Cys 140	aac Asn	att Ile	gct Ala	gcc Ala	432
gat Asp 145	atg Met	atg Met	gat Asp	gtc Val	atg Met 150	cag Gln	cta Leu	ttg Leu	gca Ala	ccc Pro 155	gat Asp	gat Asp	ggg Gly	ggg G1y	aga Arg 160	480
caa Gln	ttc Phe	gac Asp	tgg Trp	aag Lys 165	gtg Val	gtg Val	aga Arg	aga Arg	gaa Glu 170	acg Thr	agc Ser	tct Ser	ttc Phe	ttg Leu 175	acc Thr	528
gaa Glu	acc Thr	gtc val	gtg val 180	ttt Phe	ggt Gly	cgg Arg	gac Asp	caa Gln 185	gaa Glu	agg Arg	gag Glu	aaa Lys	gta Val 190	gta Val	gaa Glu	576
ttg Leu	ctg Leu	ttg Leu 195	gat Asp	tca Ser	gga Gly	tct Ser	ggt Gly 200	aac Asn	agt Ser	agc Ser	ttc Phe	tct Ser 205	gtc val	tta Leu	ccc Pro	624
ctc Leu	gtc val 210	gga Gly	atc Ile	gga Gly	ggg Gly	gtt Val 215	ggg Gly	aag Lys	acg Thr	act Thr	ctg Leu 220	gct Ala	cag Gln	ctc Leu	gtg Val	672
tac Tyr 225	aac Asn	gac Asp	aat Asn	cgt Arg	gtc Val 230	ggc Gly	aac Asn	tat Tyr	ttc Phe	cac His 235	ctc Leu	aag Lys	gtt Val	tgg Trp	gtc Val 240	720
tgt Cys	gta Val	tcc Ser	gac Asp	aat Asn 245	ttc Phe	aat Asn	gtg val	aag Lys	aga Arg 250	ctg Leu	acc Thr	aaa Lys	gag Glu	ata Ile 255	atc Ile	768
gag Glu	tct Ser	gct Ala	acc Thr 260	aag Lys	gtg val	gaa Glu	caa Gln	tct Ser 265	gac Asp	aaa Lys	ttg Leu	aac Asn	ttg Leu 270	gac Asp	acc Thr	816
ctg Leu	caa Gln	cag Gln 275	atc Ile	ctc Leu	aag Lys	gag Glu	aag Lys 280	att Ile	gct Ala	tca Ser	gag Glu	agg Arg 285	ttt Phe	ctg Leu	cta Leu	864
gtc Val	ctc Leu 290	gat Asp	gat Asp	gtg val	tgg Trp	agc Ser 295	gaa Glu	aac Asn	agg Arg	gat Asp	gac Asp 300	tgg Trp	gaa Glu	agg Arg	ctg Leu	912
tgc Cys 305	gca Ala	cca Pro	cta Leu	agg Arg	ttt Phe 310	gca Ala	gca Ala	aga Arg	ggc Gly	agc Ser 315	aag Lys	gtt val	ata Ile	gtc val	aca Thr 320	960
act Thr	cga Arg	gac Asp	aca Thr	aag Lys 325	att Ile	gcc Ala	agc Ser	atc Ile	att Ile 330	ggc Gly	aca Thr	atg Met	aag Lys	gaa Glu 335	att Ile	1008
tcg Ser	ctc Leu	gat Asp	ggt Gly 340	ctc Leu	cag Gln	gat Asp	gat Asp	gct Ala 345	tac Tyr	tgg Trp	gag Glu	ctg Leu	ttc Phe 350	aag Lys	aaa Lys	1056
tgt Cys	gca Ala	ttt Phe 355	ggt Gly	tct Ser	gtg Val	aac Asn	ccc Pro 360	cag Gln	Glu	cat His	Leu	gag G1u 365	ctc Leu	gag Glu	gtt Val	1104

atc Ile	ggt Gly 370	aga Arg	aag Lys	att Ile	gct Ala	ggt Gly 375	aag Lys	ttg Leu	aag Lys	ggc Gly	tca Ser 380	ccg Pro	cta Leu	gca Ala	gca Ala	1152
aaa Lys 385	aca Thr	cta Leu	gga Gly	agc Ser	ttg Leu 390	ttg Leu	cgg Arg	ttg Leu	gat Asp	gtc Val 395	agc Ser	caa Gln	gaa Glu	cac His	tgg Trp 400	1200
aga Arg	act Thr	ata Ile	atg Met	gaa Glu 405	agt Ser	gag Glu	gta Val	tgg Trp	caa Gln 410	ctg Leu	cca Pro	caa Gln	gct Ala	gaa Glu 415	aat Asn	1248
					cta Leu											1296
ctt Leu	aga Arg	cag G1n 435	tgt Cys	ttc Phe	gct Ala	ttt Phe	tgc Cys 440	gct Ala	gtg v al	ttt Phe	cac His	aaa Lys 445	gat Asp	tat Tyr	tta Leu	1344
ttc Phe	tat Tyr 450	aaa Lys	cat His	gag Glu	ttg Leu	atc Ile 455	cag Gln	act Thr	tgg Trp	att Ile	gca Ala 460	gaa Glu	ggc Gly	ttc Phe	att Ile	1392
gca Ala 465	cat His	caa G1n	gga Gly	aac Asn	aag Lys 470	agg Arg	atg Met	gaa Glu	gat Asp	gtc Val 475	gga Gly	agc Ser	agc Ser	tac Tyr	ttc Phe 480	1440
cat His	gag Glu	ctt Leu	gtt Val	aat Asn 485	agg Arg	tct Ser	ttc Phe	ttt Phe	cag Gln 490	gaa Glu	tct Ser	cgg Arg	tgg Trp	aga Arg 495	ggg GTy	1488
cga Arg	tat Tyr	gtg Val	atg Met 500	cat His	gac Asp	ctc Leu	ata Ile	cac His 505	gat Asp	ctt Leu	gcc Ala	caa Gln	ttt Phe 510	ata Ile	tca Ser	1536
gtg Val	gga Gly	gag Glu 515	tgt Cys	cat His	agg Arg	ata Ile	gat Asp 520	gat Asp	gac Asp	aag Lys	tcc Ser	aaa Lys 525	gag Glu	acc Thr	cct Pro	1584
agt Ser	acg Thr 530	act Thr	cgt Arg	cat His	cta Leu	tca Ser 535	gta Val	gca Ala	tta Leu	act Thr	gag G1u 540	caa Gln	atg Met	aag Lys	ttg Leu	1632
gtg Val 545	gat Asp	ttt Phe	tca Ser	ggt Gly	tac Tyr 550	aat Asn	aaa Lys	ttg Leu	cgg Arg	acc Thr 555	ctt Leu	atg Met	atc Ile	aac Asn	aat Asn 560	1680
cag Gln	aga Arg	aat Asn	cag Gln	tat Tyr 565	cca Pro	tat Tyr	atg Met	act Thr	aaa Lys 570	gtc Val	aac Asn	agc Ser	tgc Cys	ttg Leu 575	ttg Leu	1728
					aaa Lys											1776
cag Gln	aag Lys	tgt Cys 595	ggc Gly	atg Met	aaa Lys	gag Glu	ttg Leu 600	cct Pro	gat Asp	att Ile	atc Ile	ggt Gly 605	gac Asp	ttg Leu	ata Ile	1824
caa G1n	ctt Leu	cgg Arg	tac Tyr	ctt Leu	gac Asp	ata Ile	tcc Ser	tac Tyr	Asn	gct Ala Page	Cys	att Ile	cag Gln	agg Arg	ttg Leu	1872

610 615

ccc Pro 625	gag Glu	tca Ser	ttg Leu	tgc Cys	gac Asp 630	ctt Leu	tac Tyr	aat Asn	ctg Leu	caa Gln 635	gca Ala	ctg Leu	agg Arg	cta Leu	tgg Trp 640	1920
ggc Gly	tgt Cys	caa Gln	tta Leu	cgg Arg 645	agt Ser	ttc Phe	cca Pro	caa Gln	ggc G1y 650	atg Met	agc Ser	aag Lys	ctg Leu	atc Ile 655	aac Asn	1968
ttg Leu	agg Arg	caa Gln	ctt Leu 660	cgt Arg	gta Val	gaa Glu	gat Asp	gag G1u 665	ata Ile	att Ile	tcc Ser	aag Lys	ata Ile 670	tat Tyr	gag Glu	2016
gtt Val	ggg Gly	aag Lys 675	ctg Leu	att Ile	tct Ser	ctg Leu	caa G1n 680	gaa Glu	ttg Leu	tct Ser	gca Ala	ttc Phe 685	aaa Lys	gtg Val	cta Leu	2064
aat Asn	aat Asn 690	cat His	gga Gly	aac Asn	aaa Lys	ctt Leu 695	gca Ala	gaa Glu	cta Leu	agt Ser	ggt Gly 700	ttg Leu	aca Thr	caa Gln	ctc Leu	2112
cgc Arg 705	agc Ser	act Thr	cta Leu	cga Arg	att Ile 710	aca Thr	aat Asn	ctt Leu	gaa Glu	aat Asn 715	gta Val	ggg G1y	agt Ser	aaa Lys	gaa Glu 720	2160
gaa Glu	gca Ala	agc Ser	aag Lys	gct Ala 725	aaa Lys	ctg Leu	cac His	agg Arg	aaa Lys 730	cag Gln	tat Tyr	ctt Leu	gaa Glu	gca Ala 735	tta Leu	2208
gag Glu	tta Leu	gag Glu	tgg Trp 740	gca Ala	gct Ala	ggc G1y	cag Gln	gtt Val 745	tcc Ser	agc Ser	ttg Leu	gag Glu	cat His 750	gag Glu	tta Leu	2256
ctt Leu	gtc val	tcg Ser 755	gag Glu	gaa Glu	gta Val	ctt Leu	tta Leu 760	ggt Gly	ctc Leu	caa Gln	cca Pro	cat His 765	cac His	ttc Phe	ctc Leu	2304
aaa Lys	agt Ser 770	ttg Leu	aca Thr	atc Ile	aga Arg	ggg G1y 775	tac Tyr	agt Ser	ggt Gly	gca Ala	aca Thr 780	gta Val	ccc Pro	agt Ser	tgg Trp	2352
														gag Glu		2400
tgt Cys	aca Thr	aga Arg	ctg Leu	gag G1u 805	ggt Gly	ctt Leu	tca Ser	tat Tyr	att Ile 810	gga Gly	caa Gln	ctg Leu	cca Pro	cat His 815	ctc Leu	2448
aag Lys	gtc val	ctt Leu	cat His 820	atg Met	aag Lys	aga Arg	atg Met	cct Pro 825	gtg Val	gtg Val	aaa Lys	caa G1n	atg Met 830	agt Ser	cat His	2496
gaa Glu	tta Leu	tgt Cys 835	ggc Gly	tgt Cys	acg Thr	aaa Lys	agc Ser 840	aag Lys	ttg Leu	ttc Phe	cct Pro	agg Arg 845	cta Leu	gaa Glu	gag Glu	2544
tta Leu	gta Val 850	ctg Leu	gag Glu	gat Asp	atg Met	cca Pro 855	aca Thr	ttg Leu	aaa Lys	gaa Glu	ttc Phe 860	ccg Pro	aat Asn	ctt Leu	gca Ala	2592
caa	ctt	cct	tgt	ctc	aag	att	att	cac		aag Page		atg	ttt	gca	gta	2640

Gln Leu Pro Cys Leu Lys Ile Ile His Met Lys Asn Met Phe Ala Val 865 870 880	
aaa cat ata ggt cgt gaa tta tat ggt gat ata gag agc aat tgt ttt Lys His Ile Gly Arg Glu Leu Tyr Gly Asp Ile Glu Ser Asn Cys Phe 885 885 885	2688
cta tca tta gaa gag ctt gtg ctg cag gac atg ctg aca ttg gag gaa Leu Ser Leu Glu Glu Leu val Leu Gln Asp Met Leu Thr Leu Glu Glu 900	2736
ctc cca aat ctt gga caa ctt cca cat ctt aag gtt att cac atg aag Leu Pro Asn Leu Gly Gln Leu Pro His Leu Lys Val Ile His Met Lys 920 925 925	2784
aac atg tct gca ctg aaa ctt ata ggt cgt gaa tta tgt gat tct aga Ass Met Ser Ala Leu Lys Leu Ile Gly Arg Glu Leu Cys Asp Ser Arg 930 940	2832
gag aaa att tgg ttt cct agg cta gaa gtg cta gtg ctg aag aac atg Glu Lys Ile Trp Phe Pro Arg Leu Glu Val Leu Val Leu Lys Asn Met 945 955 950	2880
ctg gca ctg gag gaa ctc cca agc ttg gac aac ttc cgt gtc tca aga Leu Ala Leu Glu Glu Leu Pro Ser Leu Asp Asn Phe Arg Val Ser Arg 975 975	2928
ttc ttc gca tcc agt gtc gaa gta ggc cat gga ctc ttt agt gct acg Phe Phe Ala Ser Ser Val Glu Val Gly His Gly Leu Phe Ser Ala Thr 980	2976
agg aat aaa tgg ttt cca agg ctg gaa gag cta gaa atc aag ggc atg Arg Asn Lys Trp Phe Pro Arg Leu Glu Ile Lys Gly Met 1000 1005	3024
ctg aca ttt gag gaa ctc cat tct ctt gaa aaa ctg cca tgt ctc Leu fir Phe Glu Glu Leu His Ser Leu Glu Lys Leu Pro Cys Leu 1010 1020	3069
aag gtt ttc cgc atc aag gga ttg cca gca gtg aaa aag ata ggc Lys Val Phe Arg I'le Lys Gly 1025 1030	3114
cat gga tta ttt gat tct acc tgt cag aga gag tgt ttt cca agg His Gly Leu Phe Asp Ser Thr Cys Gln Arg Glu Cys Phe Pro Arg 1049	3159
ttg gaa gat ctc gta tta agc gac atg cca gca tgg gaa gag tgg Leu Glu Asp Leu Val Leu Ser Asp Met Pro Ala Trp Glu Glu Trp 1055	3204
tcg tgg gct gaa agg gag gag tta ttt tcc tgc ttg tgt aga ctt Ser Trp Ala Glu Arg Glu Leu Phe Ser Cys Leu cys Arg Leu 1070 1080	3249
aaa att gaa caa tgc ccc aaa ctt aaa tgc ttg ctt ccc atc cct Lys Ile Glu Gln Cys Pro Lys 1085 1095	3294
cat tct ctc ata aaa ctt gaa tta tgg caa gtt ggg ctg aca gga His Ser Leu The Lys Leu Glu Leu Trp Gln Val Gly 1100 1105	3339
Page 5	

ctt cca g Leu Pro G 1115	gga tta Gly Leu	tgc aaa Cys Lys	gga Gly 1120	att Ile	70 ggt Gly	8416 gga Gly	7_1. ggt Gly	TXT agc Ser 1125		act Thr		3384
act gct t Thr Ala S 1130		tca ctc Ser Leu								aat Asn		3429
aga aat c Arg Asn L 1145	tg gga _eu Gly	gaa ggg Glu Gly	ttg Leu 1150	cta Leu	tca Ser	aac Asn	cac His	ctg Leu 1155		cat His		3474
aat gct a Asn Ala I 1160	att cgg [le Arg	ata tgg Ile Trp	gaa Glu 1165	tgt Cys	gct Ala	gaa Glu	ctg Leu	ttg Leu 1170	tgg Trp	ctg Leu		3519
gtc aag a Val Lys A 1175	agg ttt Arg Phe	aga gaa Arg Glu	ttc Phe 1180	acc Thr	acc Thr	ctt Leu	gag Glu	aac Asn 1185		tca Ser		3564
agg aac t Arg Asn C 1190	tgc ccc Cys Pro	aag ctc Lys Leu	atg Met 1195	agc Ser	atg Met	aca Thr	cag Gln	tgt Cys 1200	gag Glu	gag Glu	aat Asn	3609
gac ctc c Asp Leu L 1205	tc ctc eu Leu	ccg ccg Pro Pro	tta Leu 1210	atc Ile	aag Lys	gca Ala	cta Leu	gaa Glu 1215	ttg Leu	ggt Gly	gac Asp	3654
tgt gga a Cys Gly A 1220	aat ctt Asn Leu	ggg aaa Gly Lys	tcg Ser 1225	ctg Leu	cct Pro	gga Gly	tgc Cys	cta Leu 1230	cac His	aac Asn		3699
agc tca c Ser Ser L 1235	tt act eu Thr	cag ttg Gln Leu	gcg Ala 1240	ata Ile	tcc Ser	aat Asn	tgt Cys	cca Pro 1245		atg Met		3744
tcc ctt c Ser Leu P 1250	ca agg Pro Arg	gaa gta Glu Val	atg Met 1255	ctt Leu	cac His	ttg Leu	aag Lys	gaa Glu 1260	ctt Leu	gga G1y	act Thr	3789
gta agg a Val Arg I 1265	atc gag [le Glu	aat tgt Asn Cys	gat Asp 1270	ggg G1y	ctg Leu	gga Gly	tca Ser	ata Ile 1275	gag Glu	ggt Gly	tta Leu	3834
caa gtt c Gln Val L 1280	tc aaa Leu Lys	tca ctc Ser Leu	aag Lys 1285	aga Arg	ttg Leu	gca Ala	atc Ile	ata Ile 1290		tgt Cys		3879
agg ctt t Arg Leu L 1295	ttg cta Leu Leu	aat gaa Asn Glu	ggg Gly 1300	gat Asp	gag Glu	caa Gln	ggg Gly	gag Glu 1305		ttg Leu		3924
ctg ctt g Leu Leu G 1310	gaa tta Glu Leu	tca gta Ser Val	gat Asp 1315	aaa Lys	aca Thr	gcc Ala	cta Leu	ctt Leu 1320		ctc Leu		3969
		aca cta Thr Leu								atc Ile		4014
tgg tct c Trp Ser F 1340	cct cag Pro Gln	aaa gtg Lys Val	atg Met 1345	ttt Phe	gac Asp		gag Glu ge 6	gag Glu 1350	cag Gln	gaa Glu	ttg Leu	4059

	cac His 1355					ctc Leu 1360						ttc Phe	aga Arg		4104
	aat Asn 1370	ctc Leu	cag Gln	tcc Ser	ttg Leu	cca Pro 1375	aca Thr	gag Glu	ttg Leu	cat His	acc Thr 1380	ctt Leu	cct Pro		4149
ctc Leu	cat His 1385	gct Ala	ttg Leu	gtt Val	gta Val	agt Ser 1390	gac Asp	tgc Cys	cca Pro	cag Gln	atc Ile 1395	caa Gln	tca Ser	ctg Leu	4194
ccg Pro						aca Thr 1405									4239
	tgc Cys 1415	cac His	cca Pro	gtg Val	ctg Leu	act Thr 1420	gcg Ala	caa Gln	ctg Leu	gaa Glu	aag Lys 1425	cac His	ctg Leu		4284
	atg Met 1430	aag Lys	agc Ser	tca Ser	ggt Gly	cga Arg 1435	ttt Phe	cac His	cca Pro	gtt Val	tat Tyr 1440	gca Ala	tagg	jcaacat	4333
gagt	gagga	at go	jagaa	aggg	gag	gtggaa	iga g	gaaag	jatti	c ga	ittgc	5			4380

gagtgaggat ggagaaaggg gagtggaaga gaaagatttc gattgcc

<210> 2 <211> 1441 <212> PRT

<213> Musa acuminata

Met Ser Thr Ala Leu Val Ile Gly Gly Trp Phe Ala Gln Ser Phe Ile 1 5 10 15 Gln Thr Leu Leu Asp Lys Ala Ser Asn Cys Ala Ile Gln Gln Leu Ala 20 25 30 Arg Arg Arg Gly Leu His Asp Asp Leu Arg Arg Leu Arg Thr Ser Leu
35 40 45 Leu Arg IIe His Ala IIe Leu Asp Lys Ala Glu Thr Arg Trp Asn His 50 60 60 Lys Asn Thr Ser Leu Val Glu Leu Val Arg Gln Leu Lys Asp Ala Ala 65 70 75 80 80 Tyr Asp Ala Glu Asp Leu Leu Glu Glu Leu Glu Tyr Gln Ala Ala Lys 85 90 Gln Lys Val Glu His Arg Gly Asp Gln Ĭle Ser Asp Leu Phe Ser Phe Ser Leu Ser Thr Ala Ser Glu Trp Leu Gly Ala Asp Gly Asp Asp Ala 115 120 125 Gly Thr Arg Leu Arg Glu Ile Gln Gly Lys Leu Cys Asn Ile Ala Ala 135 140 Asp Met Met Asp Val Met Gln Leu Leu Ala Pro Asp Asp Gly Gly Arg 145 150 160 Gln Phe Asp Trp Lys Val Val Arg Arg Glu Thr Ser Ser Phe Leu Thr 165 170 175 Glu Thr Val Val Phe Gly Arg Asp Gln Glu Arg Glu Lys Val Val Glu 180 185 190 Leu Leu Leu Asp Ser Gly Ser Gly Asn Ser Ser Phe Ser Val Leu Pro 195 200 205 Leu Val Gly Ile Gly Gly Val Gly Lys Thr Thr Leu Ala Gln Leu Val 210 215 220 Tyr Asn Asp Asn Arg Val Gly Asn Tyr Phe His Leu Lys Val Trp Val

230 Cys Val Ser Asp Asn Phe Asn Val Lys Arg Leu Thr Lys Glu Ile Ile 245 250 255 Glu Ser Ala Thr Lys Val Glu Gln Ser Asp Lys Leu Asn Leu Asp Thr 260 265 270 Leu Gln Gln Ile Leu Lys Glu Lys Ile Ala Ser Glu Arg Phe Leu Leu 275 280 285 Val Leu Asp Asp Val Trp Ser Glu Asn Arg Asp Asp Trp Glu Arg Leu 290 295 300 Cys Ala Pro Leu Arg Phe Ala Ala Arg Gly Ser Lys Val Ile Val Thr 305 310 320 Thr Arg Asp Thr Lys Ile Ala Ser Ile Ile Gly Thr Met Lys Glu Ile 325 330 335 Ser Leu Asp Gly Leu Gln Asp Asp Ala Tyr Trp Glu Leu Phe Lys Lys 340 345 350 Cys Ala Phe Gly Ser Val Asn Pro Gln Glu His Leu Glu Leu Glu Val Ile Gly Arg Lys Ile Ala Gly Lys Leu Lys Gly Ser Pro Leu Ala Ala 370 380 Lys Thr Leu Gly Ser Leu Leu Arg Leu Asp Val Ser Gln Glu His Trp 385 390 395 400 Arg Thr Ile Met Glu Ser Glu Val Trp Gln Leu Pro Gln Ala Glu Asn 415 Glu Ile Leu Pro Val Leu Trp Leu Ser Tyr Gln His Leu Pro Gly His Leu Arg Gln Cys Phe Ala Phe Cys Ala Val Phe His Lys Asp Tyr Leu 435 440 445 Phe Tyr Lys His Glu Leu Ile Gln Thr Trp Ile Ala Glu Gly Phe Ile 450 460 Ala His Gln Gly Asn Lys Arg Met Glu Asp Val Gly Ser Ser Tyr Phe 465 470 475 480 His Glu Leu val Asn Arg Ser Phe Phe Gln Glu Ser Arg Trp Arg Gly
485 490 495 Arg Tyr Val Met His Asp Leu Ile His Asp Leu Ala Gln Phe Ile Ser Val Gly Glu Cys His Arg Ile Asp Asp Asp Lys Ser Lys Glu Thr Pro 515 520 525 Ser Thr Thr Arg His Leu Ser Val Ala Leu Thr Glu Gln Met Lys Leu 530 540 val Asp Phe Ser Gly Tyr Asn Lys Leu Arg Thr Leu Met Ile Asn Asn 545 550 560 661 Arg Asn Gln Tyr Pro Tyr Met Thr Lys Val Asn Ser Cys Leu Leu 570 575 575 Pro His Ser Leu Phe Lys Arg Leu Lys Arg Ile His Val Leu Val Leu 580 590 Gln Lys Cys Gly Met Lys Glu Leu Pro Asp Ile Ile Gly Asp Leu Ile 595 600 605 Gln Leu Arg Tyr Leu Asp Ile Ser Tyr Asn Ala Cys Ile Gln Arg Leu 610 615 620 Pro Glu Ser Leu Cys Asp Leu Tyr Asn Leu Gln Ala Leu Arg Leu Trp 625 630 635 640 Gly Cys Gln Leu Arg Ser Phe Pro Gln Gly Met Ser Lys Leu Ile Asn 645 650 655 Leu Arg Gln Leu Arg Val Glu Asp Glu Île Île Ser Lys Île Tyr Glu 660 665 670 Val Gly Lys Leu Ile Ser Leu Gln Glu Leu Ser Ala Phe Lys Val Leu 675 680 685 Asn Asn His Gly Asn Lys Leu Ala Glu Leu Ser Gly Leu Thr Gln Leu 690 _ _ 695 _ _ 700 _ _ _ Arg Ser Thr Leu Arg Ile Thr Asn Leu Glu Asn Val Gly Ser Lys Glu 705 710 715 720 Glu Ala Ser Lys Ala Lys Leu His Arg Lys Gln Tyr Leu Glu Ala Leu 725 730 735

7084167_1.TXT Glu Leu Glu Trp Ala Ala Gly Gln Val Ser Ser Leu Glu His Glu Leu 745 750 740 Leu Val Ser Glu Glu Val Leu Leu Gly Leu Gln Pro His His Phe Leu 755 760 765 Lys Ser Leu Thr Ile Arg Gly Tyr Ser Gly Ala Thr Val Pro Ser Trp 770 775 780 Leu Asp Val Lys Met Leu Pro Asn Leu Gly Thr Leu Lys Leu Glu Asn 790 Cys Thr Arg Leu Glu Gly Leu Ser Tyr Ile Gly Gln Leu Pro His Leu 805 810 Lys Val Leu His Met Lys Arg Met Pro Val Val Lys Gln Met Ser His 820 825 830 Glu Leu Cys Gly Cys Thr Lys Ser Lys Leu Phe Pro Arg Leu Glu Glu 835 840 845 Leu Val Leu Glu Asp Met Pro Thr Leu Lys Glu Phe Pro Asn Leu Ala 855 Gln Leu Pro Cys Leu Lys Île Île His Met Lys Asn Met Phe Ala Val 865 870 875 Lys His Ile Gly Arg Glu Leu Tyr Gly Asp Ile Glu Ser Asn Cys Phe Leu Ser Leu Glu Glu Glu Leu Val Leu Gln Asp Met Leu Thr Leu Glu Glu 900 905 910 Leu Pro Asn Leu Gly Gln Leu Pro His Leu Lys Val Ile His Met Lys 915 920 925 Ash Met Ser Ala Leu Lys Leu Île Gly Arg Glu Leu Cys Asp Ser Arg 930 935 940 Glu Lys Ile Trp Phe Pro Arg Leu Glu Val Leu Val Leu Lys Asn Met 945 950 955 960 Leu Ala Leu Glu Glu Leu Pro Ser Leu Asp Asn Phe Arg Val Ser Arg 965 970 975 Phe Phe Ala Ser Ser Val Glu Val Gly His Gly Leu Phe Ser Ala Thr 980 985 990 Arg Asn Lys Trp Phe Pro Arg Leu Glu Glu Leu Glu Ile Lys Gly Met Leu Thr Phe Glu Glu Leu His Ser Leu Glu Lys Leu Pro Cys Leu Lys 1010 1015 1020 Val Phe Arg Ile Lys Gly Leu Pro Ala Val Lys Lys Ile Gly His Gly 1025 1030 1035 1040 Leu Phe Asp Ser Thr Cys Gln Arg Glu Cys Phe Pro Arg Leu Glu Asp 1045 1050 1055 Leu Val Leu Ser Asp Met Pro Ala Trp Glu Glu Trp Ser Trp Ala Glu
1060 1065 1070 Arg Glu Glu Leu Phe Ser Cys Leu Cys Arg Leu Lys Ile Glu Gln Cys 1075 1080 1085 Pro Lys Leu Lys Cys Leu Leu Pro Ile Pro His Ser Leu Ile Lys Leu 1090 1095 1100 Glu Leu Trp Gln Val Gly Leu Thr Gly Leu Pro Gly Leu Cys Lys Gly 1105 1110 1115 1120 Ile Gly Gly Ser Ser Thr Arg Thr Ala Ser Leu Ser Leu His 1125 1130 1135 Ile Ile Lys Cys Pro Asn Leu Arg Asn Leu Gly Glu Gly Leu Leu Ser 1140 1145 1150 Asn His Leu Pro His Ile Asn Ala Ile Arg Ile Trp Glu Cys Ala Glu 1155 1160 1165 Leu Leu Trp Leu Pro Val Lys Arg Phe Arg Glu Phe Thr Thr Leu Glu 1170 1180 Asn Leu Ser Ile Arg Asn Cys Pro Lys Leu Met Ser Met Thr Gln Cys 1185 1190 1200Glu Glu Asn Asp Leu Leu Leu Pro Pro Leu Ile Lys Ala Leu Glu Leu 1205 1210 1215 Gly Asp Cys Gly Asn Leu Gly Lys Ser Leu Pro Gly Cys Leu His Asn 1220 1225 1230

Leu Ser Ser Leu Thr Gln Leu Ala Ile Ser Asn Cys Pro Tyr Met Val

1235 1240 1245

Ser Leu Pro Arg Glu Val Met Leu His Leu Lys Glu Leu Gly Thr Val 1250 1255 1260 Arg Ile Glu Asn Cys Asp Gly Leu Gly Ser Ile Glu Gly Leu Gln Val 1265 1270 1280 1280 Leu Lys Ser Leu Lys Arg Leu Ala Ile Ile Gly Cys Pro Arg Leu Leu 1285 1290 1295 Leu Asn Glu Gly Asp Glu Gln Gly Glu Val Leu Ser Leu Leu Glu Leu 1305 Ser Val Asp Lys Thr Ala Leu Leu Lys Leu Ser Leu Ile Lys Asn Thr 1320 1325 Leu Pro Phe Ile His Ser Leu Arg Ile Ile Trp Ser Pro Gln Lys Val 1335 1330 1340 Met Phe Asp Leu Glu Glu Gln Glu Leu Val His Ser Leu Thr Ala Leu 1345 1350 1355 Arg Arg Leu Glu Phe Phe Arg Cys Lys Asn Leu Gln Ser Leu Pro Thr 1365 1370 1375 Glu Leu His Thr Leu Pro Ser Leu His Ala Leu Val Val Ser Asp Cys 1380 1385 1390 Pro Gln Ile Gln Ser Leu Pro Glu Lys Gly Leu Pro Thr Leu Leu Thr 1395 1400 1405 Asp Leu Gly Phe Asp His Cys His Pro Val Leu Thr Ala Gln Leu Glu 1410 1415 1420 Lys His Leu Ala Glu Met Lys Ser Ser Gly Arg Phe His Pro Val Tyr 1425 1430 1435 1440 Ala

<210> 3 <211> 3699 <212> DNA

<212> DNA <213> Musa acuminata

<220> <221> CDS <222> (1)..(3696)

<222> (1)..(3696) <400> 3

atg oct gat gtc aca cca cag gca gcg gcg gtg ttc tcc ctg gtg aat 48 Met Ala Asp Val Thr Pro Gln Ala Ala Ala Val Phe Ser Leu Val Asn 1 5 10

gaa atc ttt aac cgg tcc atc aat ttg atc gtc gcg gaa ctc cgg ttg Glu Ile Phe Asn Arg Ser Ile Asn Leu Ile Val Ala Glu Leu Arg Leu 20 25 30

cag ttg aat gcg aga gcc gag ctg aac aat ctg cag aga aca cta ttg Gln Leu Asn Ala Arg Ala Glu Leu Asn Asn Leu Gln Arg Thr Leu Leu 35 40 45

agg act cac tct ctg ctc gag gag gca aag gcg agg cgg atg act gac Arg Thr His Ser Leu Leu Glu Glu Ala Lys Ala Arg Arg Met Thr Asp 50 60

aag tet etc gtg etg tgg etg atg gag etc aag gaa tgg gec tae gac Lys Ser Leu Val Leu Trp Leu Met Glu Leu Lys Glu Trp Ala Tyr Asp 65 70 80

gcc gac gac atc ctc gac gag tac gag gcc gca gca atc cga ctg aag 2 Ala Asp Asp Ile Leu Asp Glu Tyr Glu Ala Ala Ala Ile Arg Leu Lys 85 90

240

								ctt	atc	167_ gat Asp	cat	gtg				336
gtt val	cca Pro	tta Leu 115	gcg Ala	cac His	aaa Lys	gta Val	gca Ala 120	gac Asp	atc Ile	agg Arg	aaa Lys	agg Arg 125	ttg Leu	aac Asn	ggg G1y	384
										gcg Ala						432
ccg Pro 145	ctt Leu	gat Asp	tcc Ser	acg Thr	aaa Lys 150	aga Arg	ggt Gly	gtg Val	acc Thr	act Thr 155	tct Ser	ctt Leu	ctg Leu	act Thr	gaa Glu 160	480
tct Ser	tgt Cys	att Ile	gtc Val	ggg Gly 165	cga Arg	gct Ala	caa Gln	gat Asp	aag Lys 170	gag Glu	aat Asn	ttg Leu	att Ile	cgg Arg 175	ttg Leu	528
ctg Leu	ttg Leu	gag Glu	ccc Pro 180	agc Ser	gat Asp	ggg Gly	gcg Ala	gtt Val 185	cct Pro	gtt Val	gtt Val	cct Pro	ata Ile 190	gtt Val	gga Gly	576
tta Leu	gga Gly	ggg Gly 195	gca Ala	ggg Gly	aag Lys	acg Thr	act Thr 200	ctg Leu	tct Ser	cag Gln	ctt Leu	atc Ile 205	ttt Phe	aat Asn	gac Asp	624
aag Lys	aga Arg 210	gtg Val	gag Glu	gag Glu	cat His	ttc Phe 215	cca Pro	ttg Leu	aga Arg	atg Met	tgg Trp 220	gtg Val	tgt Cys	gtg Val	tct Ser	672
gac Asp 225	gat Asp	ttt Phe	gat Asp	gtg Val	aag Lys 230	aga Arg	att Ile	act Thr	aga Arg	gag G1u 235	atc Ile	aca Thr	gag Glu	tac Tyr	gcc Ala 240	720
										ttg Leu						768
aat Asn	ctg Leu	aaa Lys	gag G1u 260	gag Glu	ata Ile	agg Arg	ggg Gly	acg Thr 265	aca Thr	ttt Phe	ttg Leu	ctt Leu	gtg Val 270	ctg Leu	gat Asp	816
										gaa Glu						864
tta Leu	gat Asp 290	gcc Ala	gga Gly	gga Gly	cgg Arg	gga G1y 295	agc Ser	gtg val	gtc Val	att Ile	gtg Val 300	acg Thr	aca Thr	cag Gln	agc Ser	912
aaa Lys 305	aag Lys	gtc val	gcc Ala	gat Asp	gtc val 310	acc Thr	ggc Gly	acg Thr	atg Met	gag Glu 315	cca Pro	tac Tyr	gtt Val	ctc Leu	gag Glu 320	960
gag Glu	tta Leu	acg Thr	gag Glu	gat Asp 325	gac Asp	agt Ser	tgg Trp	tca Ser	ctc Leu 330	atc Ile	gag Glu	agt Ser	cac His	tcc Ser 335	ttc Phe	1008

1056

agg Arg	aag Lys	ata 11e 355	gcc Ala	aag Lys	aag Lys	atc Ile	agt Ser 360	ggc GTy	cta Leu	cct Pro	tac Tyr	gga G1y 365	gca Ala	aca Thr	gca Ala	1104
atg Met	ggg G1y 370	aga Arg	tat Tyr	cta Leu	aga Arg	tct Ser 375	aag Lys	cac His	gga G1y	gaa Glu	agc Ser 380	agc Ser	tgg Trp	aga Arg	gaa Glu	1152
gtc Val 385	ttg Leu	gaa Glu	act Thr	gag Glu	act Thr 390	tgg Trp	gag Glu	atg Met	cca Pro	ccg Pro 395	gct Ala	gca Ala	agt Ser	gat Asp	gtg Val 400	1200
tta Leu	tcc Ser	gct Ala	cta Leu	agg Arg 405	aga Arg	agt Ser	tac Tyr	gac Asp	aat Asn 410	cta Leu	ccc Pro	cct Pro	cag Gln	ctg Leu 415	aag Lys	1248
ctc Leu	tgt Cys	ttt Phe	gcc Ala 420	ttc Phe	tgt Cys	gct Ala	ctg L eu	ttt Phe 425	aca Thr	aag Lys	ggc Gly	tac Tyr	agg Arg 430	ttt Phe	cga Arg	1296
aag Lys	gat Asp	aca Thr 435	ctg Leu	atc Ile	cac His	atg Met	tgg Trp 440	ata Ile	gct Ala	caa Gln	aat Asn	ttg Leu 445	att Ile	caa Gln	tca Ser	1344
aca Thr	gag Glu 450	tcg Ser	aaa Lys	aga Arg	tcg Ser	gag G1u 455	gac Asp	atg Met	gca Ala	gaa Glu	gaa G1u 460	tgc Cys	ttt Phe	gat Asp	gat Asp	1392
ttg Leu 465	gtg Val	tgc Cys	aga Arg	ttc Phe	ttc Phe 470	ttt Phe	cgg Arg	tac Tyr	tcc Ser	tgg Trp 475	ggc Gly	aac Asn	tat Tyr	gtg val	atg Met 480	1440
aat Asn	gac Asp	tca Ser	gtc Val	cat His 485	gac Asp	ctc Leu	gct Ala	cga Arg	tgg Trp 490	gtt val	tca Ser	ttg Leu	gat Asp	gaa Glu 495	tat Tyr	1488
ttt Phe	cga Arg	gca Ala	gat Asp 500	gaa Glu	gac Asp	tca Ser	cca Pro	ttg Leu 505	cat His	att Ile	tca Ser	aag Lys	cca Pro 510	att Ile	cgt Arg	1536
		tca Ser 515														1584
aac Asn	act Thr 530	ggt Gly	gga Gly	gat Asp	gct Ala	gtc Val 535	aat Asn	ccg Pro	ctc Leu	agc Ser	agt Ser 540	ttg Leu	cgc Arg	act Thr	ctc Leu	1632
ctt Leu 545	ttc Phe	tta Leu	ggc GTy	caa Gln	tct Ser 550	gag Glu	ttc Phe	cgg Arg	tcg Ser	tat Tyr 555	cat His	ctt Leu	ctt Leu	gat Asp	aga Arg 560	1680
		agg Arg														1728
gtc Val	ata Ile	aga Arg	aat Asn 580	ttg Leu	cct Pro	tct Ser	tcg Ser	gtt Val 585	gga Gly	aat Asn	ctg Leu	aaa Lys	cat His 590	ctg Leu	cgt Arg	1776
tac Tyr	ctg Leu	ggc G1y	ctg Leu	tct Ser	aat Asn	acg Thr	aga Arg	att Ile	Gln	agg Arg age	Leu	ccg Pro	gag Glu	tct Ser	gta Val	1824

aca Thr	cgt Arg 610	ctt Leu	tgc Cys	ctc Leu	ctt Leu	cag Gln 615	aca Thr	ttg Leu	cta Leu	cta Leu	gag G1u 620	ggc Gly	tgt Cys	gaa Glu	ctg Leu	1872
tgc Cys 625	agg Arg	tta Leu	cca Pro	aga Arg	agc Ser 630	atg Met	agc Ser	agg Arg	ctc Leu	gtc Val 635	aaa Lys	ctg Leu	agg Arg	cag Gln	ctc Leu 640	1920
aaa Lys	gca Ala	aat Asn	cca Pro	gat Asp 645	gta Val	att Ile	gcc Ala	gac Asp	ata Ile 650	gcc Ala	aaa Lys	gtc Val	ggg Gly	aga Arg 655	ttg Leu	1968
atc Ile	gaa Glu	ctt Leu	caa G1n 660	gag Glu	ctg Leu	aaa Lys	gcc Ala	tat Tyr 665	aat Asn	gtt Val	gac Asp	aag Lys	aaa Lys 670	aaa Lys	gga Gly	2016
cat His	ggg Gly	att Ile 675	gca Ala	gag Glu	cta Leu	agt Ser	gca Ala 680	atg Met	aat Asn	cag Gln	ctt Leu	cac His 685	ggt Gly	gat Asp	ctt Leu	2064
tcc Ser	att Ile 690	aga Arg	aac Asn	ctt Leu	caa Gln	aat Asn 695	gta Val	gag Glu	aaa Lys	acg Thr	cga Arg 700	gag Glu	tct Ser	cgg Arg	aag Lys	2112
gcg Ala 705	agg Arg	ttg Leu	gac Asp	gag Glu	aaa Lys 710	cag Gln	aag Lys	ctt Leu	aag Lys	ctc Leu 715	ttg Leu	gat Asp	ctg Leu	cga Arg	tgg Trp 720	2160
gct Ala	gac Asp	ggt Gly	agg Arg	ggt G1y 725	gcc Ala	gga Gly	gaa Glu	tgt Cys	gat Asp 730	cgt Arg	gac Asp	agg Arg	aaa Lys	gtt val 735	ctt Leu	2208
											ttg Leu					2256
tac Tyr	gga Gly	ggc G1y 755	act Thr	tca Ser	tct Ser	ccg Pro	agt Ser 760	tgg Trp	atg Met	acg Thr	gat Asp	cag Gln 765	tat Tyr	ctg Leu	ccc Pro	2304
aac Asn	atg Met 770	gaa Glu	acg Thr	att Ile	cgc Arg	ctg Leu 775	cgt Arg	agc Ser	tgc Cys	gca Ala	agg Arg 780	ttg Leu	acg Thr	gaa Glu	ctc Leu	2352
cca Pro 785	tgt Cys	ctc Leu	ggt Gly	cag Gln	ctg Leu 790	cat His	atc Ile	ctt Leu	aga Arg	cat His 795	ttg Leu	cac His	atc Ile	gat Asp	ggg Gly 800	2400
atg Met	tcc Ser	caa Gln	gtg Val	aga Arg 805	caa Gln	att Ile	aat Asn	ctg Leu	caa Gln 810	ttt Phe	tat Tyr	ggc Gly	acc Thr	gga Gly 815	gaa Glu	2448
											ata Ile					2496
agt Ser	ctg Leu	gag Glu 835	gaa Glu	tgg Trp	tcg Ser	gaa Glu	cca Pro 840	cgg Arg	aga Arg	aac Asn	tgt Cys	tgc Cys 845	tac Tyr	ttc Phe	cct Pro	2544
cgc	ctc	cat	aaa	ctg	ctg	atc	gag	gat		ccc age	agg 13	ctc	agg	aat	ctg	2592

Arg	Leu 850	нis	Lys	Leu	Leu	11e 855	Glu	Asp		Pro		Leu	Arg	Asn	Leu	
ccc Pro 865	tcc Ser	ctc Leu	cca Pro	cca Pro	aca Thr 870	ctg Leu	gaa Glu	gaa Glu	cta Leu	agg Arg 875	ata Ile	tca Ser	aga Arg	aca Thr	gga Gly 880	2640
cta Leu	gtt Val	gat Asp	ctt Leu	cca Pro 885	gga Gly	ttc Phe	cat His	gga Gly	aac Asn 890	ggt Gly	gat Asp	gtg Val	acg Thr	acg Thr 895	aat Asn	2688
gtt Val	tcc Ser	ctt Leu	tct Ser 900	tct Ser	ttg Leu	cat His	gtt Val	tcg Ser 905	gag Glu	tgt Cys	cga Arg	gaa Glu	ctg Leu 910	Arg	tcc Ser	2736
cta Leu	agc Ser	gaa Glu 915	gga Gly	ttg Leu	ttg Leu	cag Gln	cac His 920	aac Asn	ctc Leu	gtc Val	gcc Ala	ctc Leu 925	aag Lys	aca Thr	gcg Ala	2784
gca Ala	ttt Phe 930	acc Thr	gat Asp	tgt Cys	gat Asp	tct Ser 935	ctt Leu	gag Glu	ttt Phe	ttg Leu	ccg Pro 940	Ala	gaa Glu	gga G1y	ttc Phe	2832
aga Arg 945	aca Thr	gcc Ala	att Ile	tca Ser	ctt Leu 950	gaa Glu	tca Ser	ttg Leu	ata Ile	atg Met 955	act Thr	aat Asn	tgt Cys	cca Pro	ctg Leu 960	2880
cct Pro	tgc Cys	agt Ser	ttt Phe	ctt Leu 965	ttg Leu	cct Pro	tcc Ser	tct Ser	ctc Leu 970	gag Glu	cat His	cta Leu	aag Lys	ttg Leu 975	cag Gln	2928
cca Pro	tgc Cys	ctc Leu	tat Tyr 980	cca Pro	aac Asn	aac Asn	aat Asn	gag G1u 985	gat Asp	tca Ser	ctg Leu	tca Ser	aca Thr 990	tgc Cys	ttc Phe	2976
gag Glu	aac Asn	ctc Leu 995	aca Thr	tct Ser	ctt Leu	tcc Ser	ttc Phe 1000	tt Le	g ga u As	c ato	c aa e Ly	a ga s As 10	t t p C		ca aat ro Asn	3024
ctg Leu	tca Ser 1010	tca Ser	ttt Phe	t cca e Pro	ccg Pro	1 gg1 5 Gly 10	c co / Pi L5	ct c ro L	ta t eu C	gt c		ta eu 020	tca Ser	gca Ala	ctc Leu	3069
	cat His 102	Lei	g tco u Ser	cto Lei	gto val	2 aa1 As1 10	ı C	gc c ys G	ag a In A	gg c rg L	eu G	aa 1n 035	tct Ser			3114
	cag Gln 1040	Ā٦a			tco Ser		ιĞ		gc t er L		hr I		cag Gln			3159
	cgc Arg 105!	cto Lei	acc Thr	ato Met	tca Ser	cac His 106	a a s	gt t er L	tg g eu V	tt g al G	ag g lu v 1	tg a1 065	aat Asn			3204
	gat Asp 1070	Thr	a ggg	g cto / Lei	gcg Ala	Phe 107	e A:	at a sn I	tc a le T	ct c hr A	rg T	gg rp 080	atg Met			3249
	aca Thr 108	Gly	t gad / Asp	gac S Asp	ggc Gly	tte Lei 109	J Me	tg c et L	tc a eu A	ga ca rg H	is A	ga rg 095	gca Ala			3294

	tca Ser 1100	ttt Phe	ttc Phe	ggg GTy	gga Gly	ctt Leu 1105	ctg L eu	caa	cac	7_1. ctc Leu	acc	ttc Phe	ctc Leu	cag Gln	3339
	cta Leu 1115	aag Lys	atc Ile	tgc Cys	cag Gln	tgt Cys 1120	cca Pro	caa G1n	ctc Leu	gta Val	acc Thr 1125	ttc Phe	acc Thr		3384
	gag Glu 1130	gaa Glu	gag Glu	aag Lys	tgg Trp	aga Arg 1135	aac Asn	ctt Leu	act Thr	tct Ser	ctt Leu 1140	caa Gln	att Ile	ctg Leu	3429
cac His	atc Ile 1145	gtt Val	gat Asp	tgt Cys	cca Pro	aac Asn 1150	ctg Leu	gag Glu	gta Val	ctg Leu	cct Pro 1155	gca Ala	aac Asn	ttg Leu	3474
caa Gln	agc Ser 1160	ctc Leu	tgc Cys	tcc Ser	ctc Leu	agc Ser 1165	acc Thr	ttg Leu	tac Tyr	atc Ile	gtc Val 1170	aga Arg	tgc Cys	cca Pro	3519
aga Arg	atc Ile 1175	cat His	gcg Ala	ttt Phe	cct Pro	ccc Pro 1180	gga Gly	ggt Gly	gtc val	agc Ser	atg Met 1185	tcc Ser	ctg Leu	gca Ala	3564
	ttg Leu 1190	gtc val	atc Ile	cat His	gaa Glu	tgc Cys 1195	cct Pro	cag Gln	ctg Leu	tgt Cys	cag Gln 1200	cga Arg	tgt Cys	gat Asp	3609
cca Pro	ccg Pro 1205	gga Gly	ggt Gly	gat Asp	gat Asp	tgg Trp 1210	ccc Pro	tta Leu	ata Ile	gct Ala	aat Asn 1215	gta val	cca Pro	aga Arg	3654
ata Ile	tgt Cys 1220	ctt Leu	gga Gly	agg Arg	act Thr	cat His 1225	cca Pro	tgt Cys	cgc Arg	tgt Cys	agc Ser 1230	acc Thr	acc Thr	tga	3699

<210> 4 <211> 1232 <212> PRT

<213> Musa acuminata

Ser Cys Ile Val Gly Arg Ala Gln Asp Lys Glu Asn Leu Ile Arg Leu 165 170 175 Leu Leu Glu Pro Ser Asp Gly Ala Val Pro Val Val Pro Ile Val Gly Leu Gly Gly Ala Gly Lys Thr Thr Leu Ser Gln Leu Ile Phe Asn Asp 195 200 205 Lys Arg Val Glu Glu His Phe Pro Leu Arg Met Trp Val Cys Val Ser 210 215 220 220 Asp Asp Phe Asp Val Lys Arg Ile Thr Arg Glu Ile Thr Glu Tyr Ala 225 230 235 240 Thr Asn Gly Arg Phe Met Asp Leu Thr Asn Leu Asn Met Leu Gln Val 245 250 255 Asn Leu Lys Glu Glu Ile Arg Gly Thr Thr Phe Leu Leu Val Leu Asp 265 270 Asp Val Trp Asn Glu Asp Pro Val Lys Trp Glu Ser Leu Leu Ala Pro 275 280 285 Leu Asp Ala Gly Gly Arg Gly Ser Val Val Ile Val Thr Thr Gln Ser 290 295 300 Lys Lys Val Ala Asp Val Thr Gly Thr Met Glu Pro Tyr Val Leu Glu 305 310 315 320 Glu Leu Thr Glu Asp Asp Ser Trp Ser Leu Ile Glu Ser His Ser Phe 325 330 335 Arg Glu Ala Ser Cys Ser Ser Thr Asn Pro Arg Met Glu Glu Ile Gly 340 345 350 Arg Lys Ile Ala Lys Lys Ile Ser Gly Leu Pro Tyr Gly Ala Thr Ala 355 360 365 Met Gly Arg Tyr Leu Arg Ser Lys His Gly Glu Ser Ser Trp Arg Glu 370 375 380 Val Leu Glu Thr Glu Thr Trp Glu Met Pro Pro Ala Ala Ser Asp Val 385 390 395 400 Leu Ser Ala Leu Arg Arg Ser Tyr Asp Asn Leu Pro Pro Gln Leu Lys 405 410 415 Leu Cys Phe Ala Phe Cys Ala Leu Phe Thr Lys Gly Tyr Arg Phe Arg Lys Asp Thr Leu Ile His Met Trp Ile Ala Gln Asn Leu Ile Gln Ser 435 440 445 Thr Glu Ser Lys Arg Ser Glu Asp Met Ala Glu Glu Cys Phe Asp Asp 450 460 Leu Val Cys Arg Phe Phe Phe Arg Tyr Ser Trp Gly Asn Tyr Val Met 465 470 475 480 Asn Asp Ser Val His Asp Leu Ala Arg Trp Val Ser Leu Asp Glu Tyr 485 490 495 Phe Arg Ala Asp Glu Asp Ser Pro Leu His Ile Ser Lys Pro Ile Arg His Leu Ser Trp Cys Ser Glu Arg Ile Thr Asn Val Leu Glu Asp Asn 515 520 Asn Thr GTy GTy Asp Ala Val Asn Pro Leu Ser Ser Leu Arg Thr Leu 530 _ _ 535 . 540 Leu Phe Leu Gly Gln Ser Glu Phe Arg Ser Tyr His Leu Leu Asp Arg 545 550 556 Met Phe Arg Met Leu Ser Arg Ile Arg Val Leu Asp Phe Ser Asn Cys 565 570 575 Val Ile Arg Asn Leu Pro Ser Ser Val Gly Asn Leu Lys His Leu Arg 580 585 590 Tyr Leu Gly Leu Ser Asn Thr Arg Ile Gln Arg Leu Pro Glu Ser Val 595 600 605 Thr Arg Leu Cys Leu Leu Gln Thr Leu Leu Glu Gly Cys Glu Leu 610 615 620 Cys Arg Leu Pro Arg Ser Met Ser Arg Leu Val Lys Leu Arg Gln Leu 625 630 635 640 Lys Ala Asn Pro Asp Val Ile Ala Asp Ile Ala Lys Val Gly Arg Leu 645 650 655 Ile Glu Leu Gln Glu Leu Lys Ala Tyr Asn Val Asp Lys Lys Gly

665 His Gly Ile Ala Glu Leu Ser Ala Met Asn Gln Leu His Gly Asp Leu 675 680 685 Ser Ile Arg Asn Leu Gln Asn Val Glu Lys Thr Arg Glu Ser Arg Lys 690 695 700 Ala Arg Leu Asp Glu Lys Gln Lys Leu Lys Leu Leu Asp Leu Arg Trp 705 710 715 720 Ala Asp Gly Arg Gly Ala Gly Glu Cys Asp Arg Asp Arg Lys Val Leu 725 730 735 Lys Gly Leu Arg Pro His Pro Asn Leu Arg Glu Leu Ser Ile Lys Tyr 740 745 750 Tyr Gly Gly Thr Ser Ser Pro Ser Trp Met Thr Asp Gln Tyr Leu Pro
755 760 765 Asn Met Glu Thr Ile Arg Leu Arg Ser Cys Ala Arg Leu Thr Glu Leu 770 775 780 Pro Cys Leu Gly Gln Leu His Ile Leu Arg His Leu His Ile Asp Gly 785 790 795 800 Met Ser Gln Val Arg Gln Ile Asn Leu Gln Phe Tyr Gly Thr Gly Glu 805 810 815 Val Ser Gly Phe Pro Leu Leu Glu Leu Leu Asn Ile Arg Arg Met Pro 820 825 830 Ser Leu Glu Glu Trp Ser Glu Pro Arg Arg Asn Cys Cys Tyr Phe Pro 835 840 845 Arg Leu His Lys Leu Leu Ile Glu Asp Cys Pro Arg Leu Arg Asn Leu 850 860 Pro Ser Leu Pro Pro Thr Leu Glu Glu Leu Arg Ile Ser Arg Thr Gly 865 870 875 880 Leu Val Asp Leu Pro Gly Phe His Gly Asp Gly Asp Val Thr Thr Asn 885 890 895 Val Ser Leu Ser Ser Leu His Val Ser Glu Cys Arg Glu Leu Arg Ser 900 905 910 Leu Ser Glu Gly Leu Leu Gln His Asn Leu Val Ala Leu Lys Thr Ala 915 920 925 Ala Phe Thr Asp Cys Asp Ser Leu Glu Phe Leu Pro Ala Glu Gly Phe 930 935 940 Arg Thr Ala Ile Ser Leu Glu Ser Leu Ile Met Thr Asn Cys Pro Leu 945 950 955 960 Pro Cys Ser Phe Leu Leu Pro Ser Ser Leu Glu His Leu Lys Leu Gln 965 970 975 Pro Cys Leu Tyr Pro Asn Asn Asn Glu Asp Ser Leu Ser Thr Cys Phe 980 985 990 Glu Asn Leu Thr Ser Leu Ser Phe Leu Asp Ile Lys Asp Cys Pro Asn 995 1000 1005 Leu Ser Ser Phe Pro Pro Gly Pro Leu Cys Gln Leu Ser Ala Leu Gln 1010 1015 1020 His Leu Ser Leu Val Asn Cys Gln Arg Leu Gln Ser Ile Gly Phe Gln 1025 1030 1035 Ala Leu Thr Ser Leu Glu Ser Leu Thr Ile Gln Asn Cys Pro Arg Leu 1045 1050 1055 Thr Met Ser His Ser Leu Val Glu Val Asn Asn Ser Ser Asp Thr Gly 1060 1065 1070 Leu Ala Phe Asn Ile Thr Arg Trp Met Arg Arg Arg Thr Gly Asp Asp 1075 1080 1085 Gly Leu Met Leu Arg His Arg Ala Gln Asn Asp Ser Phe Phe Gly Gly 1090 1095 1100 Leu Leu Gln His Leu Thr Phe Leu Gln Phe Leu Lys Ile Cys Gln Cys 1105 1110 1115 1120Pro Gln Leu Val Thr Phe Thr Gly Glu Glu Glu Lys Trp Arg Asn 1125 1130 1135 Leu Thr Ser Leu Gln Ile Leu His Ile Val Asp Cys Pro Asn Leu Glu 1140 1145 1150 Val Leu Pro Ala Asn Leu Gln Ser Leu Cys Ser Leu Ser Thr Leu Tyr

```
7084167 1.TXT
Ile Val Arg Cys Pro Arg Ile His Ala Phe Pro Pro Gly Gly Val Ser
                                                1180
                          1175
   1170
Met Ser Leu Ala His Leu Val Ile His Glu Cys Pro Gln Leu Cys Gln
1185
                      1190
                                           1195
Arg Cys Asp Pro Pro Gly Gly Asp Asp Trp Pro Leu Ile Ala Asm Val
                                        1210
                 1205
                                                              1215
Pro Arg Ile Cys Leu Gly Arg Thr His Pro Cys Arg Cys Ser Thr Thr Thr 1220 1225 1230
<210> 5
<211> 87
<212> PRT
<213> Artificial Sequence
<220>
<221> VARIANT
<222> (2)...(3)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (4)...(4)
<223> Xaa = Asn, His, Gln, Cys, Ser, Thr
<220>
<220>
<221> VARIANT
<222> (5)...(5)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (6)...(7)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (8)...(8)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (9)...(9)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (10)...(10)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (11)...(11)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
```

<222> (13)...(14)

<223> Xaa = any amino acid residue

```
<220>
<221> VARIANT
<222> (15)...(15)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (16)...(17)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (19)...(24)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (25)...(25)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (26)...(26)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (27)...(27)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (29)...(30)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (32)...(33)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (34)...(34)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (38)...(38)
<223> Xaa = any amino acid residue
```

<220>

```
<221> VARIANT
<222> (40)...(40)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (41)...(41)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (43)...(43)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (44)...(44)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (46)...(46)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (47)...(47)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (49)...(55)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (59)...(59)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (60)...(61)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<220>
<221> VARIANT
<222> (62)...(63)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (66)...(66)
\langle 223 \rangle Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (67)...(67)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (72)...(72)
```

<223> Xaa´ = Asp, Glu

```
<220>
<221> VARIANT
<222> (74)...(74)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (76)...(76)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (78)...(78)
<223> Xaa = Tvr. Val. Ile. Leu. Met. Phe. Trp
<220>
<221> VARIANT
<222> (80)...(81)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (83)...(83)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (84)...(84)
<223> Xaa = Arg, Lys, His
<221> VARIANT
<222> (85)...(85)
<223> Xaa = any amino acid residue
<400> 5
Ser Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Asn Xaa Xaa Xaa Xaa
                                      10
Xaa Leu Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Leu Xaa Xaa Leu Xaa 20 25 30
Xaa Xaa Leu Leu Arg Xaa His Xaa Xaa Leu Xaa Xaa Ala Xaa Xaa Arg
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ser Leu Val Xaa Xaa Xaa Xaa Xaa Leu 50 55 60
Lys Xaa Xaa Ala Tyr Asp Ala Xaa Asp Xaa Leu Xaa Glu Xaa Glu Xaa 65 70
Xaa Ala Xaa Xaa Xaa Lys Val
<210> 6
<211> 298
<212> PRT
<213> Artificial Sequence
<220>
<221> VARIANT
<222> (2)...(3)
<223> Xaa = any amino acid residue
<220>
```

<221> VARIANT

```
<222> (5)...(5)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (7)...(7)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<220>
<221> VARIANT
<222> (11)...(11)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (12)...(12)
<223> Xaa = any amino acid residue
<220>
<22U>
<221> VARIANT
<222> (13)...(14)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<220>
<221> VARIANT
<222> (17)...(17)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (19)...(19)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (20)...(20)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (22)...(22)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (23)...(24)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<220>
<221> VARIANT
<222> (25)...(25)
<223> Xaa = Asp, Glu, Arg, Lys, His
```

<220> <221> VARIANT

```
<222> (29)...(29)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (30)...(31)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (32)...(32)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (34)...(35)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (36)...(36)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (38)...(38)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (40)...(40)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (42)...(42)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (45)...(45)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (48)...(48)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (54)...(54)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (57)...(58)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
```

```
<220>
<221> VARIANT
<222> (61)...(61)
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (64)...(66)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (68)...(68)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (70)...(70)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (71)...(71)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<220>
<221> VARIANT
<222> (78)...(78)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (80)...(80)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (84)...(84)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (86)...(86)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (89)...(89)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (91)...(91)
<223> Xaa = any amino acid residue
<220>
<220>
<221> VARIANT
<222> (94)...(95)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
```

<222> (96)...(96)

```
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<220>
<221> VARIANT
<222> (97)...(98)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (100)...(101)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (104)...(105)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (108)...(109)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (113)...(113)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (115)...(115)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (116)...(116)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (117)...(118)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (128)...(128)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (130)...(132)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (133)...(133)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (136)...(136)
```

<223> Xaa = any amino acid residue

<220>

```
<221> VARIANT
<222> (138)...(138)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (142)...(142)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (143)...(143)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (144)...(145)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<220>
<221> VARIANT
<222> (149)...(149)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (155)...(157)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (159)...(159)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (161)...(161)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (162)...(162)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (163)...(163)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (167)...(167)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (168)...(168)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (169)...(169)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
                                           Page 26
```

```
<220>
<221> VARIANT
<222> (170)...(170)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (172)...(172<u>)</u>
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (173)...(173)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (175)...(175)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (176)...(176)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (178)...(179)
<223> Xaa = any amino acid residue
<221> VARIANT
<222> (181)...(181)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (183)...(183)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (184)...(184)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (185)...(186)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (187)...(187)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (189)...(193)
<223> Xaa = any amino acid residue
```

<220> <221> VARIANT

```
<222> (194)...(194)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (195)...(198)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (199)...(199)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (200)...(200)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (202)...(202)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (209)...(209)
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (211)...(211)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (212)...(212)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (214)...(214)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (216)...(216)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (217)...(217)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
```

```
<220>
<221> VARIANT
<222> (219)...(219)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (220)...(220)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (221)...(221)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<220>
<221> VARIANT
<222> (223)...(223)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (224)...(224)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (227)...(227)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (228)...(228)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (229)...(229)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (230)...(230)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (231)...(233)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (236)...(236)
<223> Xaa = anv amino acid residue
```

```
<220>
<221> VARIANT
<222> (237)...(238)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (240)...(240)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (242)...(242)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (244)...(244)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (245)...(245)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<220>
<221> VARIANT
<222> (247)...(247)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (249)...(250)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (251)...(251)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (252)...(252)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (254)...(254)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (255)...(255)
<223> Xaa = anv amino acid residue
<220>
<220>
<221> VARIANT
<222> (257)...(258)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
```

<222> (261)...(262)

```
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (265)...(268)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (269)...(269)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (276)...(276)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (278)...(278)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (280)...(280)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (282)...(282)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (284)...(284)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (286)...(286)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (287)...(287)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (290)...(291)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (295)...(296)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (297)...(297)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<400> 6
```

```
7084167_1.TXT
Arg Xaa Xaa Thr Xaa Ser Xaa Leu Thr Glu Xaa Xaa Xaa Xaa Gly Arg
1 5 10 15
Xaa Gln Xaa Xaa Glu Xaa Xaa Xaa Leu Leu Leu Xaa Xaa Xaa Xaa Xaa Xaa 30 30
Gly Xaa Xaa Xaa Phe Xaa Val Xaa Pro Xaa Val Gly Xaa Gly Gly Xaa
35 40 45
Gly Lys Thr Thr Leu Xaa Gln Leu Xaa Xaa Asn Asp Xaa Arg Val Xaa
50 55 60
Xaa Xaa Phe Xaa Leu Xaa Xaa Trp Val Cys Val Ser Asp Xaa Phe Xaa
                      70
Val Lys Arg Xaa Thr Xaa Glu Ile Xaa Glu Xaa Ala Thr Xaa Xaa Xaa
                 85
                                        90
Xaa Xaa Asp Xaa Xaa Asn Leu Xaa Xaa Leu Gln Xaa Xaa Leu Lys Glu
             100
                                   105
Xaa Ile Xaa Xaa Xaa Xaa Phe Leu Leu Val Leu Asp Asp Val Trp Xaa
115 120 125
Glu Xaa Xaa Xaa Trp Glu Xaa Leu Xaa Ala Pro Leu Xaa Xaa Xaa
                          135
Xaa Arg Gly Ser Xaa Val Ile Val Thr Thr Xaa Xaa Xaa Lys Xaa Ala
                      150
                                            155
Xaa Xaa Xaa Gly <u>Thr</u> Met Xaa Xaa Xaa Xaa Leu Xaa Xaa Leu Xaa Xaa
                 165
                                        170
Asp Xaa Xaa Trp Xaa Leu Xaa Xaa Xaa Xaa Xaa Phe Xaa Xaa Xaa Xaa Xaa 180 185 190
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu Xaa Ile Gly Arg Lys Ile Ala
195 200 205
Xaa Lys Xaa Xaa Gly Xaa Pro Xaa Xaa Ala Xaa Xaa Gly Xaa Xaa
210 215 220
Leu Arg Xaa Xaa Xaa Xaa Xaa Xaa Xaa Trp Arg Xaa Xaa Xaa Glu Xaa
225 230 235 240
Glu Xaa Trp Xaa Xaa Pro Xaa Ala Xaa Xaa Xaa Leu Xaa Xaa Leu 245 250 255
Xaa Xaa Ser Tyr Xaa Xaa Leu Pro Xaa Xaa Leu Xaa Xaa Cys Phe Ala
260 265 270
Phe Cys Ala Xaa Phe Xaa Lys Xaa Tyr Xaa Phe Xaa Lys Xaa Xaa Leu
275 280 285
Ile Xaa Xaa Trp Ile Ala Xaa Xaa Xaa Ile
<210> 7
<211> 285
<212> PRT
<213> Artificial Sequence
<220>
<221> VARIANT
<222> (2)...(2)
<223> xaa = any amino acid residue
<221> VARIANT
<222> (3)...(3)
<223> Xaa = Asp, Glu, Arg, Lys His
<220>
<221> VARIANT 
<222> (4)...(4)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
```

<222> (5)...(5)

```
7084167 1.TXT
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (7)...(7)
<223> Xaa = Arg, Lys, His
<220>
<220>
<221> VARIANT
<222> (8)...(8)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (10)...(10)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (13)...(13)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (16)...(16)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (17)...(17)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (18)...(19)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (21)...(21)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (22)...(22)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (23)...(23)
<223> Xaa = Arg, Lys, His
<220>
```

```
<221> VARIANT
<222> (24)...(24)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (27)...(28)
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (29)...(29)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (31)...(31)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (33)...(34)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (39)...(39)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (40)...(40)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (42)...(42)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (44)...(44)
<223> Xaa = Glv, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (45)...(45)
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (53)...(53)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (54)...(54)
<223> Xaa = any amino acid residue
```

<220>

```
<221> VARIANT
<222> (55)...(55)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (57)...(58)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (61)...(61)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (63)...(63)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (65)...(65)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (68)...(68)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (70)...(71)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (72)...(72)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (74)...(74)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (75)...(75)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (78)...(78)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (80)...(80)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (81)...(81)
<223> Xaa = any amino acid residue
                                           Page 35
```

```
<220>
<221> VARIANT
<222> (86)...(86)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (87)...(89)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (90)...(90)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (91)...(91)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (93)...(93)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (94)...(94)
<223> Xaa = Asp, Glu, Arg, Lys, His
<221> VARIANT
<222> (96)...(96)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (97)...(97)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (100)...(100)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (103)...(103)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (108)...(108)
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (110)...(110)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (111)...(111)
```

```
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (113)...(114)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (115)...(115)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (116)...(116)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (118)...(119)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (120)...(120)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (125)...(125)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (126)...(126)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (127)...(127)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (130)...(130)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (131)...(131)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (132)...(132)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (134)...(134)
<223> Xaa = any amino acid residue
```

<220>

```
<221> VARIANT
<222> (136)...(136)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (139)...(139)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (142)...(144)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (145)...(145)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<220>
<221> VARIANT
<222> (147)...(147)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (148)...(148)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (151)...(151)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (153)...(154)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (157)...(157)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (159)...(159)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (160)...(160)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (162)...(162)
<223> Xaa = Asp, Glu
<220>
<221> VARIANT
<222> (164)...(164)
<223> Xaa = Asp, Glu, Arg, Lys, His
```

```
<220>
<221> VARIANT
<222> (167)...(167)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (169)...(175)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (177)...(180)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (181)...(182)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (185)...(185)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (188)...(188)
<223> Xaa = any amino acid residue
<221> VARIANT
<222> (191)...(192)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (194)...(194)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (195)...(195)
<223> Xaa = any amino acid residue
<220>
<220>
<221> VARIANT
<222> (197)...(197)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (199)...(199)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (200)...(200)
<223> Xaa = any amino acid residue
<220>
```

<221> VARIANT

```
<222> (202)...(202)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (204)...(205)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (206)...(206)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (210)...(210)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (211)...(213)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (214)...(214)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (218)...(218)
<223> Xaa = Tvr. Val. Ile. Leu. Met. Phe. Trp
<220>
<221> VARIANT
<222> (219)...(219)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (221)...(221)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (222)...(222)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (224)...(224)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (225)...(225)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (227)...(227)
```

<223> Xaa = Gly, Ser, Ala, Thr, Pro

```
<220>
<221> VARIANT
<222> (230)...(231)
<223> Xaa = anv amino acid residue
<220>
<221> VARIANT
<222> (233)...(233)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (234)...(234)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (235)...(235)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (239)...(240)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (242)...(242)
<223> Xaa = Arg, Lys, His
<220>
<221> VARIANT
<222> (243)...(243)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (246)...(246)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (247)...(247)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (248)...(248)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (250)...(250)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<220>
<221> VARIANT
<222> (251)...(251)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
```

<222> (253)...(253)

```
<223> Xaa = Arg. Lvs. His
<220>
<221> VARIANT
<222> (255)...(255)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (256)...(258)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (259)...(259)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<220>
<221> VARIANT
<222> (260)...(260)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (262)...(262)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (263)...(263)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (264)...(264)
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (265)...(267)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (270)...(270)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (273)...(273)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (275)...(275)
<223> Xaa = any amino acid residue
<220>
<221> VARIANT
<222> (276)...(276)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
```

<220>

```
<223> Xaa = Asp, Glu, Arg, Lys, His
<220>
<221> VARIANT
<222> (281)...(281)
<223> Xaa = Gly, Ser, Ala, Thr, Pro
<220>
<221> VARIANT
<222> (283)...(283)
<223> Xaa = Asp, Glu, Arg, Lys, His
<221> VARIANT
<222> (285)...(285)
<223> Xaa = Tyr, Val, Ile, Leu, Met, Phe, Trp
<400> 7
Leu Xaa Xaa Xaa Phe Xaa Xaa Leu Xaa Arg Ile Xaa Val Leu Xaa
Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Leu Pro Xaa Xaa Xaa Gly Xaa Leu
20 25 30
Xaa Xaa Leu Arg Tyr Leu Xaa Xaa Ser Xaa Asn Xaa Xaa Ile Gln Arg
                            40
                                                45
Leu Pro Glu Ser Xaa Xaa Xaa Leu Xaa Xaa Leu Gln Xaa Leu Xaa Leu
                        55
                                            60
Xaa Gly Cys Xaa Leu Xaa Xaa Xaa Pro Xaa Xaa Met Ser Xaa Leu Xaa
65
                    70
                                         75
Xaa Leu Arg Gln Leu Xaa Xaa Xaa Xaa Xaa Ile Xaa Xaa Ile Xaa
                85
                                                         95
Xaa Val Gly Xaa Leu Ile Xaa Leu Gln Glu Leu Xaa Ala Xaa Xaa Val
            100
                                105
                                                     110
Xaa Xaa Xaa Gly Xaa Xaa Xaa Ala Glu Leu Ser Xaa Xaa Xaa Gln
        115
                             120
Leu Xaa Xaa Xaa Leu Xaa Ile Xaa Asn Leu Xaa Asn Val Xaa Xaa Xaa
    130
                                             140
                        135
Xaa Glu Xaa Xaa Lys Ala Xaa Leu Xaa Xaa Lys Gln Xaa Leu Xaa Xaa
145 150 160
Leu Xaa Leu Xaa Trp Ala Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu
165 170 175
                                                         175
Xaa Xaa Xaa Xaa Xaa Xaa Val Leu Xaa Gly Leu Xaa Pro His Xaa Xaa
            180
                                                     190
                                185
Leu Xaa Xaa Leu Xaa Ile Xaa Xaa Tyr Xaa Gly Xaa Xaa Xaa Pro Ser
        195
                            200
                                                 205
Trp Xaa Xaa Xaa Xaa Leu Pro Asn Xaa Xaa Thr Xaa Xaa Leu Xaa
    210
                        215
                                             220
Xaa Cys Xaa Arg Leu Xaa Xaa Leu Xaa Xaa Xaa Gly Gln Leu Xaa Xaa
                    230
Leu Xaa Xaa Leu His Xaa Xaa Xaa Met Xaa Xaa Val Xaa Gln Xaa Xaa
                245
                                     250
Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Phe Pro Xaa Leu Glu
            260
                                265
Xaa Leu Xaa Xaa Xaa Met Pro Xaa Leu Xaa Glu Xaa
        275
                            280
```

<221> VARIANT <222> (277)...(278)